****************** (MT)

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MPsrch_pp protein - protein database search, using Smith-Waterman algorithm

Run on: Sat May 13 09:23:13 2000; MasPar time 3.41 Seconds 194.564 Million cell updates/sec

Tabular output not generated.

Title:
Description:
Perfect Score:
Sequence:

>US-09-331-631-28 (1-28) from US09331631.pep 225 1 LDPIRQQQLCQMRCQQQEKDPRQQQQCK 28

Scoring table: PAM 150 Gap 15

Searched:

Post-processing: Minimum Match 0% Listing first 45 summaries

188963 segs, 23686106 residues

Database: a-geneseq35 1:geneseqp

Statistics: Mean 20.583; Variance 75.953; scale 0.271

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1 2 3 3 4 4 4 4 4 7 7 6 5 7 7 7 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Result
225 119 117 1117 1117 1113 1111 104 96 81 81 81 81 74 74 77 77 72 72 72	Score
100.0 52.0 52.0 52.0 52.0 52.0 52.0 52.0	Query Match
28 525 525 526 666 666 666 666 919 919 919 919 1420 3131 1313 1313 1313 108 86	Length
	BB
W62841 W62831 R20181 R20181 R202828 W62828 W6282832 R102833 R112223 R112223 R112223 R112223 W14783 W	ij
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ALIGNMENTS

PR P	Dp. R H C	SO CCC	PPR PPR	RESULT ID W AC W DT 2 DE S
Qy 1 LDPIRQQCLCQMRCQQQEXDPRQQQCK 28 RESULT 2 ID W62830 standard; Protein; 625 AA. AC W62830; First entry) DT 27-0CT-1998 (first entry) DE Macadamia integrifolia antimicrobial protein. WW antimicrobial protein; infestation; control. WW Macadamia integrifolia. SMacadamia integrifolia. FI Rey Location/Qualifiers FI Peptide /note= "signal peptide" FI Protein /note= "signal peptide" FI Protein /note= "mature protein" PN W09827805-A1. PD 22-DEC-1996; AU0874. PR 20-DEC-1997; AU0874. PR 20-DEC-1997; AU0874.	Query Match 100.0%; Score 225; DB 1; Length 28; Best Local Similarity 100.0%; Pred. No. 1.69e-15; Matches 28; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 1 LDPIRQQQLCQMRCQQQEKDPRQQQQCK 28	WPI; 98-377279/32. Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals Claim 1; Page 66; 96pp; English. The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian animals. Sequence 28 AA;	W9827805-Al. 02-JUL-1998. 22-DEC-1997. AU0874. 20-DEC-1996; AU-004275. (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY. Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;	ULT 1 W62841 standard; Protein; 28 AA. W62841; W62841; 27-OCT-1998 (first entry) Stenocarpus sinuatus antimicrobial protein. antimicrobial protein; infestation; control.

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Best Local Similarity 65.4%;
Matches 17; Conservative
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Best Local 9
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26-DEC-1991.
07-JUN-1991; G00914.
11-JUN-1990; GB-013016.
(MRSC) MARS UK LTD.
Spencer ME, Hodge R, Deakin E
WPD: 92-024418/03.
N-PSDB; 020377.
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02-JUL-1998.
22-DEC-1997; A
20-DEC-1996; A
                           expression vectors
Claim 4; Fig 2; 59pp; English.
The inventors claim a 67 kD and 31
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mar
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N-PSDB; V42316.
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Best Local S
Matches
                                                             02-JUL-1998.
22-DEC-1997; i
20-DEC-1996; i
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22-DEC-1997;
20-DEC-1996;
  (RETR-) CO
Bower NI,
WPI; 98-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be used animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocoa beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close homologies between the 67 kD polypeptide and the vicilins, which are seed storage proteins.
                                                                                                                                                                                                                                                                                                                   T 6
W62829 standard; Protein;
W62829;
27-OCT-1998 (first entry)
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Bower NI, Goulter KC,
WPI; 98-377279/32.
N-PSDB; V42310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel anti-microbial protein from e useful for controlling microbial in. Claim 1, Page 34-36; 96pp; English. The sequence is that of an antimicro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
                                                                                                                           WO9827805-A1.
                                                                                                                                                                       Protein
                                                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                          Macadamia integrifolia
                                                                                                                                                                                                                                                                            antimicrobial protein; infestation;
                                                                                                                                                                                                                                                                                                    Macadamia
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r NI, Goulter KC,
98-377279/32.
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Similarity 60.9%;
14; Conservative
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29. .666
/note= "mature |
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Green JL, Manners JM, Marcus
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Pred. No. 1.19e-03,
2; Mismatches 8
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infestations of plants or mamma
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s in plants and
                        Marcus
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Best Local (
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Key
cds
                                                                                                                 W62832
W62832;
27-OCT 1998 (first entry)
Gossypium hirsutum antimicrobial protein.
antimicrobial protein; infestation; control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1: Column 31-36; 24pp; English.

The mino acid sequence of the murine Ah receptor protein. The Ah receptor is a soluble protein which mediates an individuals response to variety of drugs, carcinogens and toxic agents. The gene was isolated from a mouse genomic DNA library using the probes Q99603-5. These probe were determined from the N-terminal sequence of a purified Ah receptor, which was purified from C57BL/6J mouse liver. The screen isolated a clone designated cAhl containing the full length cDNA. The 1.4 kb fragment of the murine clone was used to obtain clones covering the coding sequence of the human AH receptor (Q99602). The Ah receptors used to generate recombinant organisms useful as biomonitors for environmental pollutants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JAN-1995.
08-APR-1993; 045806.
08-APR-1993; US-045806.
(NOUN ) UNIV NORTHWESTERN & FOUND.
(WISC) WISCONSIN ALUMNI RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine Ah receptor protein.

Mouse Ah receptor; drug; carcinogen; toxic agent; probe; human; bioassay; C57BL/6J mouse liver; environmental pollutant; recombinant organism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or man Claim 1; Page 39-41; 96pp; English.

The sequence is that of an antimicrobial protein which can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid sequences encoding murine and human Ah receptors used in competitive binding assays to detect environmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bradfield CA, Dol. WPI; 95-051315/07.
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                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                             QQQLCQMRCQQQEKDPRQQQ
                                                                                                                 standard;
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larity 60.0%;
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1. .2418
                                                                                                              Protein; 590
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                                                                                                                                                                                                                                                                                                                                     Score 104; DB 1; L
Pred. No. 9.42e-03;
3; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Poland
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Pred. No. 1
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L.88e-03;
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Matches 1
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Best Local S
Matches 1
                                                                                                                                                                                                                                            commercially available human testis and prostate lambda gtil cDNA libraries. The sequence is very similar to that of rat AR and in the DNA-binding domain it is identical to that of rAR DNA-binding domain. Homology comparisons with other known steroid receptors indicate that hAR is more closely related to glucocorticoid, mineralo-corticoid and progesterone receptors than to v-erb-A or t receptors for oestrogen, vitamin D and thyroid hormones.
Androgen receptor.

Androgen receptor; acidic fibroblast growth factor; aFGF; antisense; benign prostatic hyperplasia; prostate cancer; therapy.
                                               W14783 standard; Protein; W14783; 22-JUN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                           Androgen receptor and TR2 DNA binding proteins - DNA sequences and antibodies for detection and quantification methods Claim 25; Fig 3; 79pp; English.

This sequence was deduced from a cDNA clone isolated by screen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gossypium hirsutum.
w09827805-Al.
02-JUL-1998.
22-DEC-1997: AU0874.
20-DEC-1996: AU-004275.
20-DEC-1996: AU-004275.
COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWER NI, GOULTER KC, Green JL, Manners JM, Mai
WPI: 98-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                             (ARCH-) ARCH DEV CORP.
Liao S, Chang C;
WPI; 91-178048/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hAR;
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R12223;
20-AUG-1991 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel anti-microbial protein from e.g. Macadamia integrifo useful for controlling microbial infestations of plants or Claim 1, Page 49-51, 96pp; English.

The sequence is that of an antimicrobial protein which can
                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; Q12001
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17-NOV-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human androgen
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                                                                                                                                                   62 QQQQQQQQQQQQETSPRQQQQ 83
                                                                                                                                                                           36.0%;
Local Similarity 63.6%;
es 14; Conservation
                                                                                                                              5 RQQQLCQMRCQQQEKDPRQQQQ
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Similarity 56.5%;
13; Conservative
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                                                                                                                                                                                                                                     918 AA;
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US-438775.
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1.61e+00;
-ches 6;
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5.79e-02;
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plants or man
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19-OCT-1989.
13-APR-1989. U01548.
14-APR-1988: US-182646.
(UYNC-) University of North Ca.
French FS, Wilson EM, Joseph Di
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20-SEP-1995; US-004018.
(WORC-) WORCESTER FOUND B.
Zamecnik PA;
WPI; 97-202879/18.
N-PSDB; T63407.
W33628 standard; Protein; 539 AA.
W33628;
06-JUL-1998 (first entry)
Yeast transcriptional activator factor TAF-68 protein.
TAF-68; transcriptional activator factor; transactivat
                                                                                                                                                                                                                                                                         DNA encoding androgen receptor protein - useful for transforming eukaryotic hosts for protein expression and subsequent antibody prodn. Disclosure; Fig. 4; 41pp; English.
Androgen receptor protein (AR) is used to produce mono- or poly-clonal antibodies. These are used for the detection and quantification of AR in the presence of endogenous androgen, as androgen will not interfere with binding. They may be used in assays to determine and quantify cellular distribution of AR in tumour tissue, and are esp. useful for evaluating prostate cancers to determine responsiveness to androgen withdrawal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
Human
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Human androgen receptor (W14783) binds testosterone and, acting at the transcriptional level, regulates the growth of normal prostatic cells. Antisense oligonucleotides (see also T63200, T63404-05) based on an androgen receptor cDNA clone (see also
                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 89-324206/44.
N-PSDB; N91772.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T63407) can be used to prevent androgen receptor gene expression, thereby inhibiting the growth or survival of prostatic cells for the treatment of benign prostatic hyperplasia and prostate cancer
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WO9711170-A1.
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androgen receptor; ployclonal antibody; cancer
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14; Conser
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14; Conser
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63.6%;
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Pred.
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Pred.
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h DR, Lubahn
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No. 1.61e+00;
                                                                                                                                                                                                      81; DB 1; I
No. 1.61e+00;
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         transactivating
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RESULT
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CC This polypeptide comprises a 68 kDa yeast transactivating factor, CC TAF-68, that is associated with the TATA-box binding protein.

CC TAF-68 can be isolated from Saccharomyces cerevisiae cells by CC virtue of its affinity to fungal or human TATA-box binding proteins CC using chromatographic procedures. It can also be recombinantly CC produced in host cells utilising vectors carrying a TAF-47 nucleic CC acid (see V04771). The yeast TAF complex includes 9 polypeptides CC of 180, 145 (see W33632), 116, 90 (see W33634), 68, 62, 51-54, 47 CC (see W33637) and 30 kDa. At least some of these are required for CC transcription in vitro. TAFs are also essential for viability. CC TAF-68 can be used in the discovery design and development of CC antifungal agents, e.g. for treatment of Candida albicans infections CC (candidiasis). It can also be used to raise diagnostically useful SC sequence 539 AA.
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Best Local S
Matches 1
17-JUN-1998; U12689.
17-JUN-1997; US-049728.
(USGO') US GOVERNMENT:
Meltzer P, Trent JM;
WPI; 99-080946/07.
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W81025 standa
W81025;
12-MAY-1999
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AIB1; amplified in breast cancer; cancer; steroid;
coactivator; SCR; estrogen; ER; estrogen dependent
breast cancer; lung cancer; colon cancer; prostate
                                                                                                                                   WO9857982-A2.
23-DEC-1998.
                                                                                                                                                                                                            Region
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26-JUN-1996;
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Similarity 44.0%;
11; Conservative
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396. .715;
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Pred. No.
8; Misma
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CC The AIB1 gene is amplified and over-expressed in certain cancers in particular breast cancer and steroid hormone responsive cancers. The AIB1 polypeptide can be used to identify compounds which inhibit condicates aberrantly proliferating cells, thus detection of increased expression of the AIB1 gene can be used to diagnose cancer or a predisposition towards developing cancer. Compounds which inhibit expression of the AIB1 or compounds which inhibit interaction of AIB1 or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 33.8%;
Best Local Similarity 47.8%;
Matches 11; Conservative
                                                                                                                                                                           claim.1; Fig 1; 67pp; French.

This amino acid sequence was deduced from the nucleotide sequence of a chicken nov gene clone isolated from a gene bank prepared from chicken embryonic fibroblasts screened with a tumour-derived probe. The only hydrophobic region occurs within the putative signal peptide suggesting that the protein is secreted. The protein also contains the consensus motif of proteins which bind to insulin-like growth factors. It is known that the human ICFII gene is overexpressed in some Wilm's tumours and a similar deregulation of ICFII expression could be involved in nephroblastoma development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JUN-1992; F00589.
25-JUN-1991; FR-007807.
(CNRS ) CENT NAT RECH SCI.
Martinerie C, Perbal B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1234 FROORVAMMMQQQQQQQQQQQQQQ 1256
                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleotide sequences hybridising to regions of chicken useful as probes for detecting complementary sequences evaluate development and/or differentiation of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       avian nephroblastoma; avian myeloblastoma virus; insulin-like growth factor; Wilm's tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77 14
R3 1599;
R31599;
24-MAY-1993 (first entry)
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WPI; 93-036377/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; Q36031.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 IRQQQLCQMRCQQQEKDPRQQQQ 26
                                                                                                                                                            deduced nov protein sequence contains
                                                                                                                residues.
351 AA;
                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= signal_peptide
/note= "only hydrophobic region of protein
56. .63
/label= IGF-binding_site_motif
/note= "corresponds to GCGCCXXC consensus"
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                                       32.9%;
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Pred. No.
6; Misma
              Score 74; DI
Pred. No. 7.7
7; Mismatch
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                      Mismatches
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7.25e+00;
                                                                                                                                                            39
                                                                                                                                                            (non-clustered)
                                                             Length 351;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IGF binding site;
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                 Gaps
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Search completed: Sat May
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Best Local :
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30-OCT-1997;
30-OCT-1996;
                                                                                                                                                                                     expression product produces a polyglutamine sequence from Gln-166 to Gln-188. In the normal gene there are 15-25 CAG repeats but in SCA2 patients this number is increased to 35-100. Paptides encoded by nucleic acid fragments (DNA or RNA) containing sequences from the SCA2 associated gene, antibodies recognising the peptides and antisense nucleic acids hybridising with the nucleic acid fragments can be used for the investigation and diagnosis of SCA2. They can also be used for the treatment of SCA2 by antisense therapy or gene therapy.
                                                                                                                                                                                                                                                                                                                     This is the protein sequence of a gene causative of spinocerebellar ataxia type 2 (SCA2), a neurodegenerative disease. The gene associated with SCA2, has a tri-nucleotide (CAG) repeat region which in the
                                                                                                                                                                                                                                                                                                                                                                                                 Sanpei K, Tsuji S;
WPI; 98-277215/24.
WPSDB; V30270.
Nucleic acid fragments associated with spinocerebellar ataxia type
contain increased number of CAG repeat region compared to normal
                                                                                                                                                                                                                                                                                                                                                                                      gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SRLS-) SRL INC.
Sanpei K, Tsuji
WPI; 98-272215/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Spinocerebellar ataxia type 2 (SCA2) disease associated protein. Spinocerebellar ataxia type 2; SCA2; gene therapy; antisense therapy; CAG repeat; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W60213 standard; Protein; | 1313 W60213;
                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9818920-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Region
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                                                                            232
                                                                                                                                                                                                                                                                                                                                                                   laim 1; Pages 13-22; 38pp; Japanese
                                                                                                                           Match 32.9%;
Local Similarity 50.0%;
                                            1 LDPIRQQQLCQMRCQQQEKDPRQQQQ
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                                                                                                             13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP-304059.
                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers 166. .188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
13 09:23:21 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Glutamine rich region; this region increased in SCA2 patients"
                                                                                                                              Score 74;
Pred. No.
                                                                                                                            Pred.
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                                              26
                                                                                                            Mismatches
                                                                                                                              7.25e+00;
                                                                                                                                           DB 1;
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                                                                                                                                             Length 1313;
                                                                                                             Indels
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